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(54) Title: TRANSPORTERS AND ION CHANNELS

(57) Abstract: The invention provides human transporters and ion channels (TRICH) and polynucleotides which identify and encode TRICH. The invention also provides expression vectors, host cells, antibodies, agonists, and antagonists. The invention also provides methods for diagnosing, treating, or preventing disorders associated with aberrant expression of TRICH.





and allogenic responses in pigs, validating the idea of channel blockers as safe and efficacious immunosuppressants (Cahalan, M.D. and K.G. Chandy (1997) Curr. Opin. Biotechnol. 8:749-756).

The discovery of new transporters and ion channels and the polynucleotides encoding them satisfies a need in the art by providing new compositions which are useful in the diagnosis, prevention, and treatment of transport, neurological, muscle, immunological, and cell proliferative disorders, and in the assessment of the effects of exogenous compounds on the expression of nucleic acid and amino acid sequences of transporters and ion channels.

#### SUMMARY OF THE INVENTION

The invention features purified polypeptides, transporters and ion channels, referred to collectively as "TRICH" and individually as "TRICH-1," "TRICH-2," "TRICH-3," "TRICH-4," "TRICH-6," "TRICH-6," "TRICH-7," "TRICH-8," "TRICH-9," "TRICH-10," "TRICH-11," "TRICH-11," "TRICH-12," "TRICH-13," "TRICH-14," "TRICH-15," "TRICH-16," "TRICH-17," "TRICH-18," "TRICH-19," "TRICH-20," "TRICH-21," "TRICH-22," "TRICH-23," "TRICH-24," "TRICH-25," "TRICH-26," and "TRICH-27." In one aspect, the invention provides an isolated polypeptide selected from the group consisting of a) a polypeptide comprising an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, b) a naturally occurring polypeptide comprising an amino acid sequence at least 90% identical to an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, c) a biologically active fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27. In one alternative, the invention provides an isolated polypeptide comprising the amino acid sequence of SEQ ID NO:1-27.

The invention further provides an isolated polynucleotide encoding a polypeptide selected from the group consisting of a) a polypeptide comprising an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, b) a naturally occurring polypeptide comprising an amino acid sequence at least 90% identical to an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, c) a biologically active fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, and d) an immunogenic fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27. In one alternative, the polynucleotide encodes a polypeptide selected from the group consisting of SEQ ID NO:1-27. In another alternative, the polynucleotide is selected from the group consisting of SEQ ID NO:1-27. In another alternative, the polynucleotide is selected from the group consisting of SEQ ID NO:28-54.

Additionally, the invention provides a recombinant polynucleotide comprising a promoter

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The invention also provides a method for screening a compound for effectiveness as an agonist of a polypeptide selected from the group consisting of a) a polypeptide comprising an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, b) a naturally occurring polypeptide comprising an amino acid sequence at least 90% identical to an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, c) a biologically active fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, and d) an immunogenic fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27. The method comprises a) exposing a sample comprising the polypeptide to a compound, and b) detecting agonist activity in the sample. In one alternative, the invention provides a composition comprising an agonist compound identified by the method and a pharmaceutically acceptable excipient. In another alternative, the invention provides a method of treating a disease or condition associated with decreased expression of functional TRICH, comprising administering to a patient in need of such treatment the composition.

Additionally, the invention provides a method for screening a compound for effectiveness as an antagonist of a polypeptide selected from the group consisting of a) a polypeptide comprising an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, b) a naturally occurring polypeptide comprising an amino acid sequence at least 90% identical to an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, c) a biologically active fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, and d) an immunogenic fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27. The method comprises a) exposing a sample comprising the polypeptide to a compound, and b) detecting antagonist activity in the sample. In one alternative, the invention provides a composition comprising an antagonist compound identified by the method and a pharmaceutically acceptable excipient. In another alternative, the invention provides a method of treating a disease or condition associated with overexpression of functional TRICH, comprising administering to a patient in need of such treatment the composition.

The invention further provides a method of screening for a compound that specifically binds to a polypeptide selected from the group consisting of a) a polypeptide comprising an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, b) a naturally occurring polypeptide comprising an amino acid sequence at least 90% identical to an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, c) a biologically active fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, and d) an immunogenic fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27. The method comprises a) combining the polypeptide with at least one test compound

under suitable conditions, and b) detecting binding of the polypeptide to the test compound, thereby identifying a compound that specifically binds to the polypeptide.

The invention further provides a method of screening for a compound that modulates the activity of a polypeptide selected from the group consisting of a) a polypeptide comprising an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, b) a naturally occurring polypeptide comprising an amino acid sequence at least 90% identical to an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, c) a biologically active fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27, and d) an immunogenic fragment of a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1-27. The method comprises a) combining the polypeptide with at least one test compound under conditions permissive for the activity of the polypeptide, b) assessing the activity of the polypeptide in the presence of the test compound, and c) comparing the activity of the polypeptide in the absence of the test compound with the activity of the polypeptide in the presence of the test compound is indicative of a compound that modulates the activity of the polypeptide.

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The invention further provides a method for screening a compound for effectiveness in altering expression of a target polynucleotide, wherein said target polynucleotide comprises a sequence selected from the group consisting of SEQ ID NO:28-54, the method comprising a) exposing a sample comprising the target polynucleotide to a compound, and b) detecting altered expression of the target polynucleotide.

The invention further provides a method for assessing toxicity of a test compound, said method comprising a) treating a biological sample containing nucleic acids with the test compound; b) hybridizing the nucleic acids of the treated biological sample with a probe comprising at least 20 contiguous nucleotides of a polynucleotide selected from the group consisting of i) a polynucleotide comprising a polynucleotide sequence selected from the group consisting of SEQ ID NO:28-54, ii) a naturally occurring polynucleotide comprising a polynucleotide sequence at least 90% identical to a polynucleotide sequence selected from the group consisting of SEQ ID NO:28-54, iii) a polynucleotide having a sequence complementary to i), iv) a polynucleotide complementary to the polynucleotide of ii), and v) an RNA equivalent of i)-iv). Hybridization occurs under conditions whereby a specific hybridization complex is formed between said probe and a target polynucleotide in the biological sample, said target polynucleotide selected from the group consisting of i) a polynucleotide comprising a polynucleotide sequence selected from the group consisting of SEQ ID NO:28-54, ii) a naturally occurring polynucleotide comprising a polynucleotide sequence at least 90% identical to a polynucleotide sequence selected from the group consisting of SEQ ID NO:28-54, iii) a polynucleotide

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Trp Pro Gly Lys Ile Gly Ala Phe Val Ser Ile Thr Met Gln Asn

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Ile Gly Ala Met Ser Ser Tyr Leu Phe Ile Ile Lys Tyr Glu Leu
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                                    145
Pro Glu Val Ile Arg Ala Phe Met Gly Leu Glu Glu Thr Ser Arg
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                                    160
                                                         165
Glu Trp Tyr Leu Asn Gly Asn Tyr Leu Ile Ile Phe Val Ser Val
                170
                                    175
Gly Ile Ile Leu Pro Leu Ser Leu Leu Lys Asn Leu Gly Tyr Leu
                185
                                    190
Gly Tyr Thr Ser Gly Phe Ser Leu Thr Cys Met Val Phe Phe Val
                200
                                    205
Ser Val Val Ile Tyr Lys Lys Phe Gln Ile Pro Cys Pro Leu Pro
                215
                                    220
                                                         225
Glu Asn Gln Ala Lys Gly Ser Leu His Asp Ser Gly Val Glu Tyr
                230
                                    235
Glu Ala His Ser Asp Asp Lys Cys Glu Pro Lys Tyr Phe Val Phe
                245
                                    250
Asn Ser Gln Thr Ala Tyr Ala Ile Pro Ile Leu Val Phe Ala Phe
                260
                                    265
Val Cys His Pro Glu Val Leu Pro Ile Tyr Ser Glu Leu Lys Asp
                                    280
Arg Ser Arg Arg Lys Met Gln Thr Val Ser Asn Ile Ser Ile Thr
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                                    295
                                                         300
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                                    310
Thr Phe Tyr Gly Arg Val Glu Asp Glu Leu Leu His Ala Tyr Ser
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Val Leu Val Ala Val Thr Leu Thr Val Pro Ile Val Leu Phe Pro
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                                    355
Val Arg Thr Ser Val Ile Thr Leu Leu Phe Pro Lys Arg Pro Phe
                365
                                    370
Ser Trp Ile Arg His Phe Leu Ile Ala Ala Val Leu Ile Ala Leu
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                                    385
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                395
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                                    535
Asp Tyr Ile Gln Ala Val Thr Ser Tyr Leu Ala Pro Pro Ile Thr
                545
                                     550
                                                         555
Ala Leu Phe Leu Leu Ala Ile Phe Cys Lys Arg Val Thr Glu Pro
                560
                                     565
Gly Ala Phe Trp Gly Leu Val Phe Gly Leu Gly Val Gly Leu Leu
                575
                                     580
Arg Met Ile Leu Glu Phe Ser Tyr Pro Ala Pro Ala Cys Gly Glu
                590
                                    595
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Phe Ala Ile Leu Cys Gly Leu Thr Ala Ile Val Ile Val Ile
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                                     625
                                                         630
Leu Thr Arg Leu Thr Trp Trp Thr Arg Asn Cys Pro Leu Ser Glu
                635
                                     640
Leu Glu Lys Glu Ala His Glu Ser Thr Pro Glu Ile Ser Glu Arg
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Pro Ala Gly Glu Cys Pro Ala Gly Gly Ala Ala Glu Asn Ser
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                                     670
Ser Leu Gly Gln Glu Gln Pro Glu Ala Pro Ser Arg Ser Trp Gly
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                                     685
Lys Leu Leu Trp Ser Trp Phe Cys Gly Leu Ser Gly Thr Pro Glu
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                                    700
                                                         705
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                                     715
                                                         720
Thr Ser Ile Glu Glu Pro Leu Trp Arg His Val Cys Asn Ile
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Arg Gly Phe Ser Gln Ser Ile Thr Pro Ser Ser Gly Gly Ser Gly
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                                     85
Ile Pro Glu Leu Lys Thr Met Leu Ala Gly Val Ile Leu Glu Asp
                                    100
Tyr Leu Asp Ile Lys Asn Phe Gly Ala Lys Val Val Gly Leu Ser
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Ala Thr Lys Pro Gly Arg Ser Gly Lys Glu Ser Val Thr Glu Pro Trp Ala Arg Val Pro Gly Ala Leu Gly Val Ala Ala Arg Gln Met His Pro Lys Ser Ile Ile Thr Phe Arg Glu Ile Asn Gly Glu Tyr Thr Gly Ala Val Asp Phe Pro Arg Leu Gly Val Arg Ala Ser Glu Glu Thr Ala Leu Arg Glu Leu Lys Met Ser Lys Glu Leu Ala Ala Met Gly Pro Gly Ala Ser Gly Asp Gly Val Arg Thr Glu Thr Ala Pro His Ile Ala Leu Asp Ser Arg Val Gly Leu His Ala Tyr Asp Ile Ser Val Val Val Ile Tyr Phe Val Phe Val Ile Ala Val Gly Ile Trp Ser Ser Ile Arg Ala Ser Arg Gly Thr Ile Gly Gly Tyr Phe Leu Ala Gly Arg Ser Met Ser Trp Trp Pro Ile Gly Ala Ser Leu Met Ser Ser Asn Val Gly Ser Gly Leu Phe Ile Gly Leu Ala Gly Thr Gly Ala Ala Gly Gly Leu Ala Val Gly Gly Phe Glu Trp Asn Ala Thr Trp Leu Leu Leu Ala Leu Gly Trp Val Phe Val Pro Val Tyr Ile Ala Ala Gly Val Val Thr Met Pro Gln Tyr Leu Lys Lys Arg Phe Gly Gly Gln Arg Ile Gln Val Tyr Met Ser Val Leu Ser Leu Ile Leu Tyr Ile Phe Thr Lys Ile Ser Thr Asp Ile Phe Ser Gly Ala Leu Phe Ile Gln Met Ala Leu Gly Trp Asn Leu Tyr Leu Ser Thr Gly Ile Leu Leu Val Val Thr Ala Val Tyr Thr Ile Ala Gly Gly Leu Met Ala Val Ile Tyr Thr Asp Ala Leu Gln Thr Val Ile Met Val Gly Gly Ala Leu Val Leu Met Phe Leu Gly Phe Gln Asp Val Gly Trp Tyr Pro Gly Leu Glu Gln Arg Tyr Arg Gln Ala Ile Pro Asn Val Thr Val Pro Asn Thr Thr Cys His Leu Pro Arg Pro Asp Ala Phe His Ile Leu Arg Asp Pro Val Ser Gly Asp Ile Pro Trp Pro Gly Leu Ile Phe Gly Leu Thr Val Leu Ala Thr Trp Cys Trp Cys Thr Asp Gln Val Ile Val Gln Arg Ser Leu Ser Ala Lys Ser Leu Ser His Ala Lys Gly Gly Ser Val Leu Gly Gly Tyr Leu Lys Ile Leu Pro Met Phe Phe Ile Val Met Pro Gly Met Ile Ser Arg Ala Leu Phe Pro Asp Glu Val Gly Cys Val Asp Pro Asp Val Cys Gln Arg Ile Cys Gly Ala Arg Val Gly Cys Ser Asn Ile Ala Tyr Pro Lys Leu Val Met Ala Leu Met Pro Val Gly Leu Arg Gly Leu Met Ile Ala Val Ile Met Ala Ala Leu Met Ser Ser Leu Thr Ser Ile Phe Asn Ser Ser Ser Thr Leu Phe Thr Ile Asp